

## SEQUENCE LISTING

&lt;110&gt; Smith, Kelli E.

Weinshank, Richard L.

&lt;120&gt; DNA Encoding A Human Receptor (hpl5a) And Uses Thereof

&lt;130&gt; 55180

&lt;140&gt; 09/179,798

&lt;141&gt; 1998-10-27

&lt;160&gt; 16

&lt;170&gt; PatentIn Ver. 2.1

&lt;210&gt; 1

&lt;211&gt; 1311

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 1

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&lt;210&gt; 2

&lt;211&gt; 396

Applicants: Kelli E. Smith and  
Richard Weinshank  
Serial No: Not Yet Known  
Filed: Herewith  
**Exhibit 1**

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 2

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Met Trp Asn Ser Ser Asp Ala Asn Phe Ser Cys Tyr His Glu Ser Val
  1                      5                      10                      15

Leu Gly Tyr Arg Tyr Val Ala Val Ser Trp Gly Val Val Val Ala Val
          20                      25                      30

Thr Gly Thr Val Gly Asn Val Leu Thr Leu Leu Ala Leu Ala Ile Gln
      35                      40                      45

Pro Lys Leu Arg Thr Arg Phe Asn Leu Leu Ile Ala Asn Leu Thr Leu
      50                      55                      60

Ala Asp Leu Leu Tyr Cys Thr Leu Leu Gln Pro Phe Ser Val Asp Thr
      65                      70                      75                      80

Tyr Leu His Leu His Trp Arg Thr Gly Ala Thr Phe Cys Arg Val Phe
          85                      90                      95

Gly Leu Leu Leu Phe Ala Ser Asn Ser Val Ser Ile Leu Thr Leu Cys
          100                      105                      110

Leu Ile Ala Leu Gly Arg Tyr Leu Leu Ile Ala His Pro Lys Leu Phe
          115                      120                      125

Pro Gln Val Phe Ser Ala Lys Gly Ile Val Leu Ala Leu Val Ser Thr
          130                      135                      140

Trp Val Val Gly Val Ala Ser Phe Ala Pro Leu Trp Pro Ile Tyr Ile
      145                      150                      155                      160

Leu Val Pro Val Val Cys Thr Cys Ser Phe Asp Arg Ile Arg Gly Arg
          165                      170                      175

Pro Tyr Thr Thr Ile Leu Met Gly Ile Tyr Phe Val Leu Gly Leu Ser
          180                      185                      190

Ser Val Gly Ile Phe Tyr Cys Leu Ile His Arg Gln Val Lys Arg Ala
          195                      200                      205

Ala Gln Ala Leu Asp Gln Tyr Lys Leu Arg Gln Ala Ser Ile His Ser
          210                      215                      220

Asn His Val Ala Arg Thr Asp Glu Ala Met Pro Gly Arg Phe Gln Glu
      225                      230                      235                      240

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Leu Asp Ser Arg Leu Ala Ser Gly Gly Pro Ser Glu Gly Ile Ser Ser  
 245 250 255  
 Glu Pro Val Ser Ala Ala Thr Thr Gln Thr Leu Glu Gly Asp Ser Ser  
 260 265 270  
 Glu Val Gly Asp Gln Ile Asn Ser Lys Arg Ala Lys Gln Met Ala Glu  
 275 280 285  
 Lys Ser Pro Pro Glu Ala Ser Ala Lys Ala Gln Pro Ile Lys Gly Ala  
 290 295 300  
 Arg Arg Ala Pro Asp Ser Ser Ser Glu Phe Gly Lys Val Thr Arg Met  
 305 310 315 320  
 Cys Phe Ala Val Phe Leu Cys Phe Ala Leu Ser Tyr Ile Pro Phe Leu  
 325 330 335  
 Leu Leu Asn Ile Leu Asp Ala Arg Val Gln Ala Pro Arg Val Val His  
 340 345 350  
 Met Leu Ala Ala Asn Leu Thr Trp Leu Asn Gly Cys Ile Asn Pro Val  
 355 360 365  
 Leu Tyr Ala Ala Met Asn Arg Gln Phe Arg Gln Ala Tyr Gly Ser Ile  
 370 375 380  
 Leu Lys Arg Gly Pro Arg Ser Phe His Arg Leu His  
 385 390 395

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<211> 45

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: primer/probe

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<213> Artificial Sequence

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<223> Description of Artificial Sequence: primer/probe

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<210> 10

<211> 46

<212> DNA

<213> Artificial Sequence

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<210> 12

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<210> 15

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<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: primer/probe

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<210> 16

<211> 25

<212> DNA

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<223> Description of Artificial Sequence: primer/probe

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